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<150> <151>	PCT/DK04/000914 2004-12-23	
<150> <151>	PA 2004 01843 2004-11-26	
<150> <151>	PA 2004 00586 2004-04-07	
<150> <151>	PA 2004 00096 2004-01-24	
<150> <151>	PA 2003 01940 2003-12-27	
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<211> 2033

<212> DNA

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 $<\!213\!>\,$ NM_006291.2| Homo sapiens tumor necrosis factor, alpha-induced protein 2 (TNFAIP2), mRNA

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<211> 1536

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<211> 2986

<212> DNA

 $<\!213\!>\,$ NM_000201.1| Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus receptor (ICAM1), mRNA

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<212> DNA

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<210> 19

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 $<\!213\!>\,$ NM_004850.3| Homo sapiens Rho-associated, coiled-coil containing protein kinase 2 (ROCK2), mRNA

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<211> 1556

<212> DNA

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<211> 1276

<212> DNA

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<211> 1577

<212> DNA

<213> NM_006214.2| Homo sapiens phytanoyl-CoA hydroxylase (Refsum disease) (PHYH), mRNA

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<211> 3060

<212> DNA

 $<\!213\!>\,$ NM_004739.2| Homo sapiens metastais—associated gene family, member 2 (MTA2), mRNA

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<210> 24

<211> 2407

<212> DNA

 $<\!213\!>\,$ NM_001091.1| Homo sapiens amiloride binding protein 1 (amine oxidase (copper-containing)) (ABP1), mRNA

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<211> 1094

<212> DNA

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<211> 5546

<212> DNA

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<211> 2545

<212> DNA

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<212> DNA

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<212> DNA

 $<\!213\!>$ NM_014298.3| Homo sapiens quinolinate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating)) (QPRT), mRNA

<400> 29 60 teccaecece ageetgggge etetgggage ettggteetg ageageeaae acaecageee 120 agacagetge aagteaceat ggacgetgaa ggeetggege tgetgetgee geeegteace 180 ctggcagccc tggtggacag ctggctccga gaggactgcc cagggctcaa ctacgcagcc 240 ttggtcagcg gggcaggccc ctcgcaggcg gcgctgtggg ccaaatcccc tggggtactg gcagggcage ctttcttcga tgccatattt acccaactca actgccaagt ctcctggttc 300 360 ctccccgagg gatcgaagct ggtgccggtg gccagagtgg ccgaggtccg gggccctgcc 420 cactgcctgc tgctggggga acgggtggcc ctcaacacgc tggcccgctg cagtggcatt 480 gccagtgctg ccgccgctgc agtggaggcc gccagggggg ccggctggac tgggcacgtg 540 gcaggcacga ggaagaccac gccaggcttc cggctggtgg agaagtatgg gctcctggtg 600 ggcggggccg cctcgcaccg ctacgacctg ggagggctgg tgatggtgaa ggataaccat 660 gtggtggccg ccggtggcgt ggagaaggcg gtgcgggcgg ccagacaggc ggctgacttc 720 getetgaagg tggaagtgga atgeageage etgeaggagg eegtgeagge agetgagget ggtgccgacc ttgtcctgct ggacaacttc aagccagagg agctgcaccc cacggccacc 780 840 gtgctgaagg cccagttccc gagtgtggct gtggaagcca gtgggggcat caccctggac aacctccccc agttctgcgg gccgcacata gacgtcatct ccatggggat gctgacccag 900 960 geggeeeeag eeettgattt eteeeteaag etgtttgeea aagaggtgge teeagtgeee aaaatccact agtcctaaac cggaagagga tgacaccggc catgggttaa cgtggctcct 1020 1080 caggaccete tgggtcacae atetttaggg teagtggeea atggggeaea tttggeaeta 1140 gettgagece aactetgget etgecacetg etgeteetgt gaeetgteag ggetgaette acctctgctc atctcagttt cctaatctgt aaaatgggtc taataaagga tcaaccacat 1200

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<211> 768

<212> DNA

 $<\!213\!>\,$ NM_004585.2| Homo sapiens retinoic acid receptor responder (tazarotene induced) 3 (RARRES3), mRNA

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<211> 696

<212> DNA

<213> NM_002984.1| Homo sapiens chemokine (C-C motif) ligand 4 (CCL4), mRNA

<400> 31	
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<210> 32

<211> 3338

<212> DNA

 $<\!213\!>\,$ NM_001455.2| Homo sapiens forkhead box O3A (FOXO3A), transcript variant 1, mRNA

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<210> 33

<211> 2646

<212> DNA

 $<\!213\!>\,$ NM_152873.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 4, mRNA

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<211> 817

<212> DNA

 $<\!213\!>\,$ NM_002038.2| Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 1, mRNA

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<210> 35

<211> 1172

<212> DNA

<213> NM_001565.1| Homo sapiens chemokine (C-X-C motif) ligand 10 (CXCL10), mRNA

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<213> NM_000043.3 Homo sapiens tumor necrosis factor recep	otor superfamily,
member 6 (TNFRSF6), transcript variant 1, mRNA	
<400> 37	

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120

180

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<211> 1600

<212> DNA

 $<\!213\!>\,$ NM_001953.2| Homo sapiens endothelial cell growth factor 1 (platelet-derived) (ECGF1), mRNA

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<210> 39

<211> 931

<212> DNA

<213> NM_005138.1| Homo sapiens SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA

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<210> 40

<211> 1216

<212> DNA

<213> $NM_006419.1$ | Homo sapiens chemokine (C-X-C motif) ligand 13 (B-cell chemoattractant) (CXCL13), mRNA

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<211> 738

<212> DNA

 $<\!213\!>\,$ NM_006433.2| Homo sapiens granulysin (GNLY), transcript variant NKG5, mRNA

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<211> 1579

<212> DNA

<213> NM_001767.2| Homo sapiens CD2 antigen (p50), sheep red blood cell receptor (CD2), mRNA

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<211> 3738

<212> DNA

<213> NM_006275.4| Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA

<400> 43 60 ctggcgcgcg cgcgccat tgtgtggctg gactcggccg cccctgtggt gtgaggcgcg 120 tgttcgggct cttgccgtcc ccgcacccgc accgcggtta ctggcttgcg gtccgccgtt 180 cgacaaccag cccttgggtc cccgcccgcc acggacatgc cgcgcgtcta cataggacgc 240 ctgagctaca acgtccggga gaaggacatc cagcgctttt tcagtggcta tggccgcctc 300 ctcgaagtag acctcaaaaa tgggtacggc ttcgtggagt tcgaggactc ccgcgacgcc 360 gacgacgccg tttacgagct gaacggcaag gagctctgcg gcgagcgcgt gatcgtagag cacgcccggg gcccgcgtcg cgatcgcgac ggctacagct acggaagccg cagtggtgga 420 480 ggtggataca gcagtcggag aacatctggc agagacaaat acggaccacc tgttcgtaca 540 gaatacaggc ttattgtaga aaatctttct agtcggtgca gttggcaaga tttaaaggat tttatgcgac aagcaggtga agtaacctat gcggatgccc acaaggaacg aacaaatgag 600 ggtgtaattg agtttcgctc ctactctgac atgaagcgtg ctttggacaa actggatggc 660 acagaaataa atggcagaaa tattaggctt attgaagata agccacgcac aagccatagg 720 780 cgatcttact ctggaagcag atccaggtct cgatctagaa gacggtcacg aagtaggagt

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<211> 2033

<212> DNA

<213> NM_003212.1| Homo sapiens teratocarcinoma-derived growth factor 1 $\,$ (TDGF1), mRNA

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<212> DNA

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<211> 3349

<212> DNA

 $<\!\!213\!\!>\,$ NM_004602.1| Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T4, mRNA

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<212> DNA

 $<\!213\!>\,$ NM_021246.2| Homo sapiens lymphocyte antigen 6 complex, locus G6D (LY6G6D), transcript variant 1, mRNA

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<210> 52

<211> 3248

<212> DNA

<213> NM_007236.3| Homo sapiens calcium binding protein P22 (CHP), mRNA

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<211> 3098

<212> DNA

 $<\!213\!>\,$ NM_003671.2| Homo sapiens CDC14 cell division cycle 14 homolog B (S. cerevisiae) (CDC14B), transcript variant 1, mRNA

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<211> 4568

<212> DNA

 $<\!213\!>\,$ NM_012408.3| Homo sapiens protein kinase C binding protein 1 (PRKCBP1), transcript variant 2, mRNA

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<212> DNA

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<210> 59

<211> 2402

<212> DNA

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<211> 2856

<212> DNA

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<211> 1655

<212> DNA

<213> NM_002164.3| Homo sapiens indoleamine-pyrrole 2,3 dioxygenase (INDO), mRNA

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<210> 62

<211> 2242

<212> DNA

<213> NM_021784.3| Homo sapiens forkhead box A2 (FOXA2), transcript variant 1, mRNA

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<210> 63

<211> 1047

<212> DNA

<213> NM_033423.2| Homo sapiens granzyme H (cathepsin G-like 2, protein h-CCPX) (GZMH), mRNA

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<211> 5243

<212> DNA

 $<\!213\!>\,$ NM_001165.3| Homo sapiens baculoviral IAP repeat-containing 3 (BIRC3), transcript variant 1, mRNA

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<211> 3850

<212> DNA

 $<\!213\!>\,$ NM_005682.4| Homo sapiens G protein-coupled receptor 56 (GPR56), transcript variant 1, mRNA

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<211> 1209

<212> DNA

<213> NM_173834.2| Homo sapiens hypothetical protein MGC21416 (MGC21416), mRNA

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<211> 722

<212> DNA

<213> NM_175617.2| Homo sapiens metallothionein 1E (functional) (MT1E), mRNA

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<210> 72

<211> 980

<212> DNA

<213> NM_003283.3| Homo sapiens troponin T1, skeletal, slow (TNNT1), mRNA

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<210> 73

<211> 2213

<212> DNA

<213> NM_004067.1| Homo sapiens chimerin (chimaerin) 2 (CHN2), mRNA

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<211> 2201

<212> DNA

<213> NM_005520.1| Homo sapiens heterogeneous nuclear ribonucleoprotein H1 (H) (HNRPH1), mRNA

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<211> 1895

<212> DNA

 $<\!\!213\!\!>$ NM_004046.4| Homo sapiens ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle (ATP5A1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA

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<212> DNA

 $<\!213\!>\,$ NM_001970.3| Homo sapiens eukaryotic translation initiation factor 5A (EIF5A), mRNA

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<212> DNA

<213> NM_005041.3| Homo sapiens perforin 1 (pore forming protein) (PRF1), mRNA

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<211> 4623

<212> DNA

<213> NM_014965.2| Homo sapiens OGT(O-Glc-NAc transferase)-interacting protein 106 KDa (OIP106), mRNA

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<210> 79

<211> 2657

<212> DNA

<213> NM_017895.6| Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 27 (DDX27), mRNA

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<210> 80

<211> 3246

<212> DNA

 $<\!213\!>$ NM_018206.3| Homo sapiens vacuolar protein sorting 35 (yeast) (VPS35), mRNA

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<210> 81

<211> 3182

<212> DNA

<213> NM_017583.3| Homo sapiens tripartite motif-containing 44 (TRIM44), mRNA

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<211> 4930

<212> DNA

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<212> DNA

<213> NM_014183.2| Homo sapiens dynein, cytoplasmic, light polypeptide 2A (DNCL2A), transcript variant 1, mRNA

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<210> 84

<211> 2100

<212> DNA

<213> NM_015907.2| Homo sapiens leucine aminopeptidase 3 (LAP3), mRNA

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<211> 1510

<212> DNA

 $<\!213\!>$ NM_018478.1| Homo sapiens chromosome 20 open reading frame 35 (C20orf35), mRNA

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<211> 3105

<212> DNA

 $<\!213\!>\,$ NM_030674.2| Homo sapiens solute carrier family 38, member 1 (SLC38A1), mRNA

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<211> 2711

<212> DNA

<213> NM_016028.4| Homo sapiens suppressor of variegation 4-20 homolog 1 (Drosophila) (SUV420H1), transcript variant 2, mRNA

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<212> DNA

 $<\!213\!>\,$ NM_022105.2| Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 1, mRNA

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<210> 94

<211> 4372

<212> DNA

 $<\!213\!>\,$ NM_014314.2| Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 (DDX58), mRNA

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<211> 2163

<212> DNA

 $<\!213\!>\,$ NM_015515.3| Homo sapiens keratin 23 (histone deacetylase inducible) (KRT23), transcript variant 1, mRNA

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<210> 96

<211> 2881

<212> DNA

 $<\!213\!>$ NM_007210.2| Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6) (GALNT6), mRNA

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<211> 1930

<212> DNA

<213> NM_020183.3| Homo sapiens aryl hydrocarbon receptor nuclear translocator-like 2 (ARNTL2), mRNA

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<211> 5730

<212> DNA

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<210> 102

<211> 2368

<212> DNA

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<211> 1672

<212> DNA

<213> NM_004503.2| Homo sapiens homeo box C6 (HOXC6), transcript variant 1, mRNA

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<213> NM_004764.2| Homo sapiens piwi-like 1 (Drosophila) (PIWIL1), mRNA

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<210> 107

<211> 2524

<212> DNA

 $<\!213\!>\,$ NM_000249.2| Homo sapiens mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), mRNA

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<212> DNA

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<212> DNA

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<211> 3262

<212> DNA

 $<\!213\!>\,$ NM_002860.2| Homo sapiens aldehyde dehydrogenase 18 family, member Al (PYCS/ALDH18A1), mRNA

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<210> 111

<211> 2899

<212> DNA

<213> NM_005655.1 \mid Homo sapiens TGFB inducible early growth response (TIEG), mRNA

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<211> 3138

<212> DNA

 $<\!213\!>\,$ NM_018223.1| Homo sapiens checkpoint with forkhead and ring finger domains (CHFR), mRNA

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<211> 2466

<212> DNA

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<211> 3919

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<211> 7401

<212> DNA

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<211> 2745

<212> DNA

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<211> 2152

<212> DNA

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<210> 120

<211> 3010

<212> DNA

<213> NM_145343.1 \mid Homo sapiens apolipoprotein L, 1 (APOL1), transcript variant 2, mRNA

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<211> 2759

<212> DNA

 $<\!213\!>\,$ NM_080796.1| Homo sapiens death associated transcription factor 1 (DATF1), transcript variant 2, mRNA

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<210> 122

<211> 781

<212> DNA

<213> NM_177953.1| Homo sapiens dynein, cytoplasmic, light polypeptide 2A (DNCL2A), transcript variant 2, mRNA

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<210> 123

<211> 841

<212> DNA

 $<\!213\!>\,$ NM_022873.1| Homo sapiens interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 3, mRNA

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<211> 4652

<212> DNA

 $<\!213\!>\,$ NM_183047.1| Homo sapiens protein kinase C binding protein 1 (PRKCBP1), transcript variant 1, mRNA

<400> 124 gtgagaacta ggagcctgtc ctccatgttt tataagtatt gacattacac agtgttaaca 60 atgcatccac agagettggc tgaagaggaa ataaaaaacag aacaggaggt ggtagagggc 120 atggatatet etaetegete caaagateet ggetetgeag agagaacage eeagaaaaga 180 240 aagttcccca gccctccaca ttcttccaat ggccactcgc cgcaggacac atcaacaagc cccattaaaa agaaaaaqaa acctqqctta ctqaacaqta acaataaqqa qcaqtcaqaa 300 360 ctaagacatg gtccgtttta ctatatgaag cagccactca ccacagaccc tgttgatgtt 420 gtaccgcagg atggacggaa tgatttctac tgctgggttt gtcaccggga aggccaagtc ctttgctgtg agctctgtcc ccgggtttat cacgctaagt gtctgagact gacatcggaa 480 ccagaggggg actggttttg tcctgaatgt gagaaaatta cagtagcaga atgcatcgag 540 acccagagta aagccatgac aatgctcacc attgaacagt tatcctacct gctcaagttt 600 gccattcaga aaatgaaaca gccagggaca gatgcattcc agaagcccgt tccattggaa 660 cagcaccctg actatgcgga atacatcttc catccaatgg acctttgtac attggaaaag 720 780 aatgcgaaaa agaaaatgta tggctgcaca gaagccttcc tggctgatgc aaagtggatt ttgcacaact gcatcattta taatggggga aatcacaaat tgacgcaaat agcgaaagta 840 900 qtcatcaaaa tctqtqaaca tqaqatqaat qaaatcqaaq tatqtccaqa atqttatcta 960 gctgcttgcc aaaaacgaga taactggttt tgtgagcctt gtagcaatcc acatcctttg gtctgggcca aactgaaggg gtttccattc tggcctgcaa aagctctaag ggataaagac 1020 1080 gggcaggtcg atgcccgatt ctttggacaa catgacaggg cctgggttcc aataaataat 1140 tgctacctca tgtctaaaga aattcctttt tctgtgaaaa agactaagag catcttcaac agtgccatgc aagagatgga ggtttacgtg gagaacatcc gcaggaagtt tggggttttt 1200 aattactctc catttaggac accctacaca cccaacagcc agtatcaaat gctgctcgat 1260 cccaccaacc ccagcgccgg cactgccaag atagacaagc aggagaaggt caagctcaac 1320 1380 tttgacatga cggcatcccc caagatcctg atgagcaagc ctgtgctgag tgggggcaca

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 $<\!\!213\!\!>\,$ NM_017452.1| Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T2, mRNA

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<211> 3506

<212> DNA

<213> NM_017453.1| Homo sapiens staufen, RNA binding protein (Drosophila) (STAU), transcript variant T3, mRNA

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<212> DNA

 $<\!\!213\!\!>\,$ NM_199169.1| Homo sapiens transmembrane, prostate androgen induced RNA (TMEPAI), transcript variant 2, mRNA

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<211> 4531

<212> DNA

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<210> 129

<211> 2692

<212> DNA

<213> NM_152871.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 2, mRNA

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<212> DNA

<213> NM_152872.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 3, mRNA

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<211> 2563

<212> DNA

 $<\!\!213\!\!>$ NM_152874.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 8, mRNA

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<211> 2445

<212> DNA

 $<\!213\!>\,$ NM_152876.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 6, mRNA

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<212> DNA

<213> NM_152877.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 7, mRNA

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<212> DNA

 $<\!213\!>\,$ NM_152875.1| Homo sapiens tumor necrosis factor receptor superfamily, member 6 (TNFRSF6), transcript variant 5, mRNA

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<211> 316

<212> DNA

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<213> NM_000251. Homo sapiens mutS...[gi:4557760]

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<211> 3239

<212> DNA

<213> NM_000534. Homo sapiens PMS1...[gi:53729349]

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<211> 2771

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<213> NM_000535. Homo sapiens PMS2...[gi:11125773]

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<212> DNA

<213> NM_000179. Homo sapiens mutS...[gi:4504190]

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